

OIPE

5/7/02

RAW SEQUENCE LISTING

DATE: 05/07/2002

PATENT APPLICATION: US/09/770,517C

TIME: 14:01:06

Input Set : A:\sequence listing.txt

Output Set: N:\CRF3\05072002\I770517C.raw

```
Does Not Comply
4 <110> APPLICANT: Mitsuhashi, Kazuya
                                                                Corrected Diskette Needed
        Yamamoto, Hiroaki
5
        Matsuyama, Akinobu
        Tokuyama, Shinji
9 <120> TITLE OF INVENTION: D-AMINOACYLASE AND GENE ENCODING THE SAME
11 <130> FILE REFERENCE: 06501-072001
13 <140> CURRENT APPLICATION NUMBER: US 09/770,517C
14 <141> CURRENT FILING DATE: 2001-01-26
16 <150> PRIOR APPLICATION NUMBER: JP 2000-019080
17 <151> PRIOR FILING DATE: 2000-01-27
19 <150> PRIOR APPLICATION NUMBER: JP 2000-150578
20 <151> PRIOR FILING DATE: 2000-05-22
22 <160> NUMBER OF SEQ ID NOS: 27
24 <170> SOFTWARE: PatentIn Ver. 2.0
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ERRORED SEQUENCES

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464 <210> SEQ ID NO: 27
     465 <211> LENGTH: 20
     466 <212> TYPE: PRT
     467 <213> ORGANISM: Hypomyces mycophilus
     469 <400> SEQUENCE: 27
     470 Val Leu Ala Asp Glu Tyr Pro Gln Ala Phe Tyr Ala Pro His Ala Tyr
                                               10
     472 Ser Arg Gly Phe
     473
E--> 474(
E--> 485
```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/770,517C

DATE: 05/07/2002 TIME: 14:01:07

Input Set : A:\sequence listing.txt
Output Set: N:\CRF3\05072002\I770517C.raw

L:157 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:161 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:172 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:176 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:187 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:191 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
L:192 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:474 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:474 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
L:485 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:27